

RESULT		8
source	1124446	
ORGANISM	AU124446	859 bp mRNA
DEFINITION	AU124446	EST
SEQUENCE	NT_2RM4	clone N12RM4000018 5', mRNA
ACCESSION	AU124446	23'-WT-206:0
VERSION	AU124446.1	EST
KEYWORDS	G1:109494946	
SOURCE	human.	
ORGANISM	Homo sapiens	
EUKARYOTA: Metazoa: Chordata: Craniata: Vertebrata: Euteleostomi: Mammalia: Eutheria: Primates: Catarrhini: Hominoidea: Homo: (bases 1 to 859)		
REFERENCE	1	
AUTHORS	Okada T., Wakanuma A., Ozawa M., Ishii S., Saito K., Yamamoto J., Nakamura Y., Nishikawa T., Nagai T., Suzuki Y., Suyano S., and Isono T.	
TITLE	Helix human cDNA project (Okada T., Wakanuma A., Ozawa M., Ishii S., Y., Nishikawa T., Nagai T., Suzuki Y., Nishikawa T., Nagai T., Suzuki T., Suyano S., and Isono T.)	
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HELIIX	Helix Research Institute	
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NOTE	human cDNA project; 5' - & 3' - end one pass sequencing; Helix Research Institute cDNA library construction, Department of Virology, Institute of Medical Science, University of Tokyo, and Helix Research Institute	
LOCATION/QUALIFIERS		
1.	'859	
	/organism="Homo sapiens"	
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	/clone_id="N12RM4"	
	/cell_type="neuronal cell line"	
	/cell_line="NT2"	
	/note="vector: pME18SFL3; mRNA from uninduced NT2 neurons precursor cells"	
USE COUNT	197 a	220 t
ALIGN	247 c	192 t
		3 others
Query Match	80.5%	Score 483; DB 10; Length 859;
Best Local Similarity	92.9%	Prog No 5, 2c, 128;

TITLE	HPI human cDNA project (Ita, T., Wakamatsu, A., Ozawa, M., Ishii, S., Saito, K., Yamamoto, J., Nakamura, Y., Nishikawa, T., Nagai, T., Suzuki, Y., Sugatani, S., Isozai, T.)
JOURNAL	Unpublished (2000)
COMMENT	Contact: 'takao.1sozai@genomix.hri.co.jp'
GENOME	Genomix Laboratory
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FEATURES	Helix human cDNA project; 5'-&-3' end one pass sequencing; Helix Research Institute, cDNA library construction, Department of Virology, Institute of Medical Science, University of Tokyo, and Helix Research Institute.
LOCATION/QUALIFIERS	source
QUALIFIER	1. 706
ORGANISM	"Homo sapiens"
DB_XREF	/taxon:9606
CLONE	"NT2RM2001157"
CLONE_ID	"NT2RM2"
CELL_TYPE	"teratocarcinoma"
CELL_LINE	"NT2"
NOTE	"vector: pME18SFL3; mRNA from undifferentiated NT2 neuroblast precursor cells"
BASE COUNT	175
ORIGIN	a 192 c 185 q 150 t 4 others

